In the Claims

Please amend page 20, line 1 as follows:

Claims What is claimed is:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

- 1. (Currently amended) NMR-based method comprising generating a NMR spectrum of a mixture comprising at least_(i) one hyperpolarised ligand, a target and optionally at least one further ligand, or (ii) a hyperpolarised target and at least one ligand and comparing said NMR spectrum with a reference spectrum of the at least one hyperpolarised ligand or the hyperpolarised target.
- 2. (Currently amended) The NMR-based method according to claim 1 comprising
 - a) hyperpolarising at least one of said a ligand or said a target,
 - b) forming a mixture by contacting <u>either</u> the at least one hyperpolarised ligand with <u>one of</u> a target <u>and or</u> a target and at least one further ligand, or the hyperpolarised target with at least one ligand,
 - c) generating a NMR spectrum of the mixture, and
 - d) comparing said NMR spectrum with a reference spectrum of the at least one hyperpolarised ligand or the hyperpolarised target.
- 3. (Currently amended) The mMethod according to elaims 1 and 2 claim 1, wherein at least one of the ligands is selected from the group consisting of proteins, glycoproteins, lipoproteins, polypeptides, glyco-polypeptides, lipopolypeptides, peptides, carbohydrates, nucleic acids or a part, a fragment or a complex thereof and small organic molecules.
- 4. (Currently amended) The mMethod according to claims 1 to 3 claim 1, wherein at least one of the ligands is a small organic molecule of less than 2000 Da.

- 5. (Currently amended) The mMethod according to elaims 1 to 4 claim 1, wherein more than one hyperpolarised ligand is used.
- 6. (Currently amended) The mMethod according to claims 1 to 5 claim 1, wherein the target is selected from the group consisting of proteins, glycoproteins, lipoproteins, nucleic acids, polypeptides, glycopolypeptides, lipopolypeptides, peptides or a part, a fragment or a complex thereof.
- 7. (Currently amended) The mMethod according to claims 1 to 6 claim 1, wherein the at least one hyperpolarised ligand or the hyperpolarised target is an isotopically enriched ligand or target.
- 8. (Currently amended) The mMethod according to elaims 1 to 7 claim 1, wherein the at least one hyperpolarised ligand or the hyperpolarised target is selectively isotopically enriched at one or more sites in the molecule.
- 9. (Currently amended) The mMethod according to claim 8 wherein the at least one hyperpolarised ligand or the hyperpolarised target is selectively isotopically enriched at one site in the molecule with ¹³C or ¹⁵N.
- 10. (Currently amended) The mMethod according to claim 9 wherein the enrichment is a ¹³C-enrichment.
- 11. (Currently amended) <u>The mMethod according to claims 1 to 10 claim 1, wherein</u> the NMR spectrum generated is a one-dimensional NMR spectrum
- 12. (Currently amended) <u>The mMethod according to elaims 1-to-11 claim 1, wherein</u> the NMR spectrum generated is generated using low flip angles.

- 13. (Currently amended) The mMethod according to elaims 1 to 12 claim 1, wherein the comparison with the reference spectrum shows a chemical shift difference, a relaxation time difference or a NOE effect difference.
- 14. (Currently amended) A method of performing an NMR-assisted drug discover comprising the step of using one or more Use of hyperpolarised ligands and/or hyperpolarised targets in NMR assisted drug discovery.
- 15. (Currently amended) <u>A method of performing a ligand competition assay comprising the step of using one or more Use of isotopically enriched hyperpolarised ligands in ligand competition assays.</u>